

Dielectron Measurements by PHENIX using the HBD

Quark Matter 2012

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For the PHENIX Collaboration

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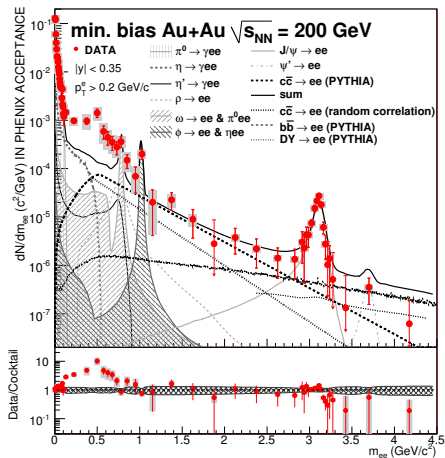
August 15, 2012



- Physics Motivation
- HBD detector
- Hadronic Cocktail
- p+p Result
- Au+Au Results

- Dileptons spectra carry abundant information about the *evolution* of the medium: They are created at all stages and do not interact strongly
- Low Mass Region
 - Modification of light vector mesons
→ Chiral Symmetry Restoration
 - Thermal photons
→ Temperature of the medium
- Intermediate and High Mass Region
 - Open heavy flavor
→ Heavy quark energy loss
 - Quarkonia
→ Quarkonium suppression
- Most striking observation is the $\times 10$ enhancement seen in the ≈ 150 MeV to 750 MeV compared to expectation from purely hadronic decays

Phys. Rev. C 81, 034911 (2010)



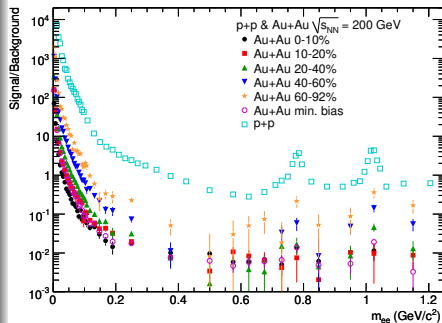
- Signal to background in PHENIX published measurement lowest in most interesting region
- Dominant source of background is combinatoric electron pairs from π^0 decays

$$\pi^0 \rightarrow \gamma e^+ e^-$$

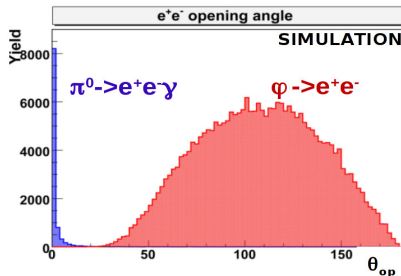
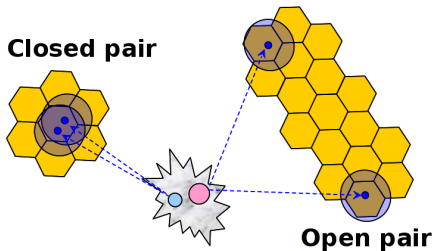
$$\pi^0 \rightarrow \gamma\gamma \rightarrow e^+ e^- e^+ e^-$$

that get separated by the magnetic field and only one leg is reconstructed

- Only way to identify such pairs is to tag them before they get separated by the magnetic field



- Identification of partially reconstructed Dalitz and early conversion:
 - Measure charge before e^+e^- pair is separated by the magnetic field (Hence the requirement of field free region and change in magnetic field configuration from previous measurements)
 - Background electrons produce twice the signal of signal electrons
 - Hadron Blind Detector (HBD)



HBD hardware

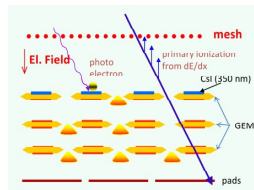
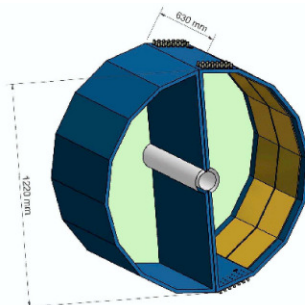
- Windowless Čerenkov Detector
- CF_4 - Radiator and Active gas
- Triple GEM signal multiplication
- CsI photo cathode, hexagonal pad readout
- 2.4% total radiation length

Hadron Blindness

- Čerenkov threshold for $\pi^\pm \approx 4 \text{ GeV}$
- Reverse bias operation to repel ionization from charged hadrons

Double rejection

- Near zero B-field up to GEMs
- Low mass pairs keep small opening angle and leave twice as much signal as single electrons



- Fit π^0 and π^\pm spectra using Hagedorn function

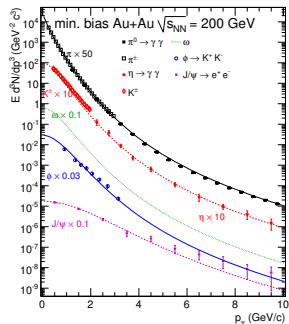
$$E \frac{d^3}{dp^3} = \frac{A}{(e^{-(ap_T + bp_T^2)} + p_T/p_0)^n}$$

- For other hadrons: Use m_T scaling for shape

$$p_T \rightarrow \sqrt{p_T^2 + (m_h^2 - m_{\pi^0}^2)}$$

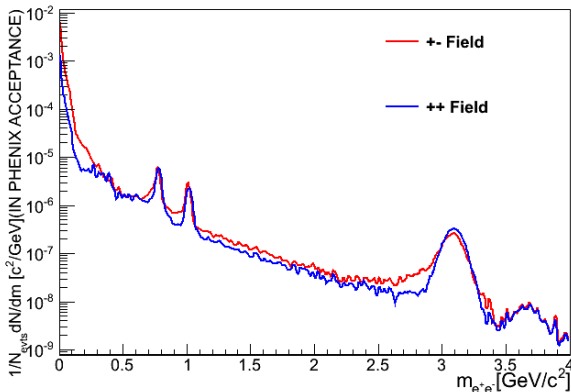
fit to measured spectra for normalization

- The fits are done independently for each species and collision centrality

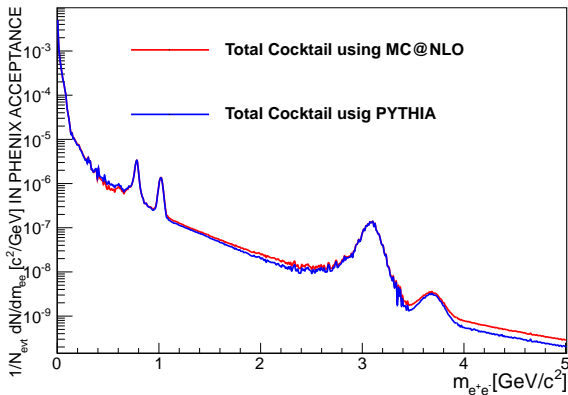


Changes from published cocktail (Phys. Rev. C 81, 034911 (2010))

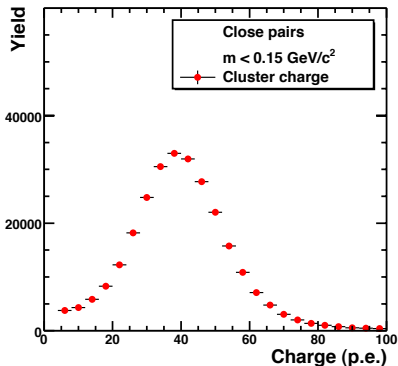
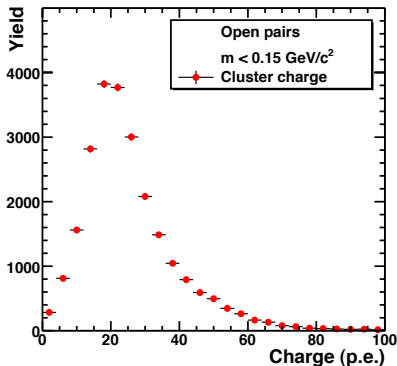
- Change in magnetic field configuration \rightarrow change in acceptance
- Open heavy flavor shape from MC@NLO as event generation
 - For p+p, charm and beauty spectra from MC@NLO are adjusted to fit data.
 - For Au+Au, an additional N_{coll} scaling is applied for each centrality bin
- J/ψ line shape extracted from full detector MC



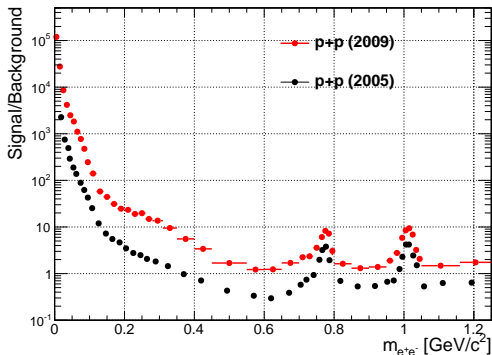
- Finite difference in pair acceptance between full field (++) and partial field (+-) configurations (HBD data was taken with (+-) field configuration)
- This difference in acceptance (shown in the mass projection above) should be accounted for in comparisons between HBD results and published phenix results.



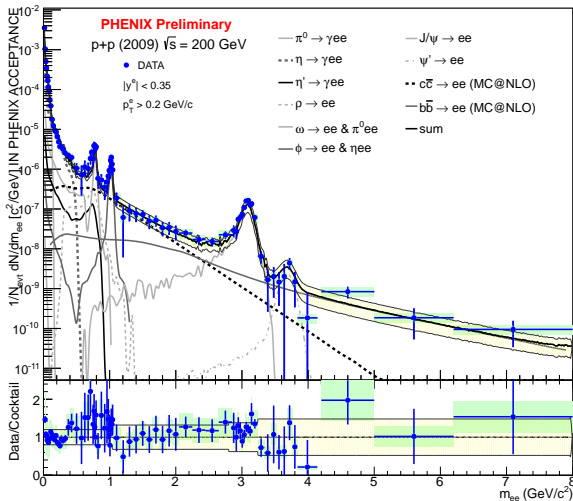
- Finite difference in total cocktail when using PYTHIA and MC@NLO as open heavy flavor event generator.
- Difference (on total cocktail) at IMR (1.2 GeV to 2.8 GeV) $\approx 16\%$ and even lower at LMR



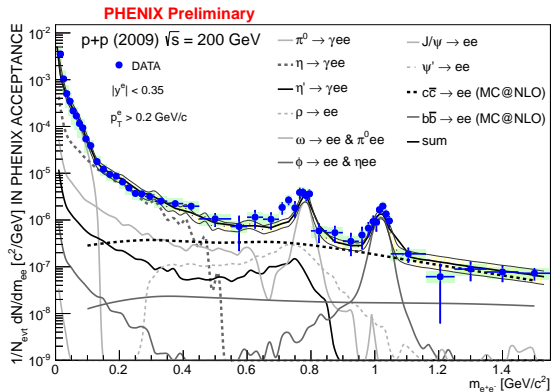
- Fully reconstructed π^0 Dalitz pairs ($m < 150 \text{ MeV}$) are split into two samples
 - Large opening angle ($> 100 \text{ mrad}$) - single signal amplitude ($\approx 20\text{pe}$)
 - Small opening angle ($< 30 \text{ mrad}$) - double signal amplitude ($\approx 40\text{pe}$)



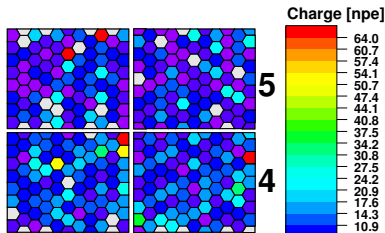
- Factor $\approx \times 5 - \times 10$ improvement in S/B in p+p
 - This improvement is achieved using the HBD just as another EID detector
 - More should be possible in p+p by using double rejection cut, but this is not the limiting systematic uncertainty in p+p results



- Excellent agreement between data and cocktail.



- Excellent agreement between data and cocktail.
- Baseline for Au+Au analysis, as well as simpler environment to test understanding of the HBD detector subsystem



- High Occupancy $\approx 100\%$ in central events mostly due to scintillation in CF_4
- Mean single signal $\approx 20\text{pe}$ over 3 pads
- Mean background $\approx 10\text{pe}$ per pad highly fluctuating response to scintillation and curlers
- \Rightarrow High fake id rate

Event By Event BG Subtr.

- **Stream A** : Mean charge per pad calculated module by module in each event
- **Stream B** : Median charge per pad calculated for each track near the projection point

- Both **Stream A** and **Stream B** use track projection based cluster charge reconstruction and achieve compatible efficiency and rejection

Au+Au Results

- The results shown now for Au+Au are only for 20-40%, 40-60% and 60-92%

- Two independent analysis were performed to increase confidence in results
- The methods were substantially different in how they handled the HBD reconstruction, electron identification, single/double rejection and correlated background subtraction
- In both analyses, the combinatorial background is subtracted using mixed events. The difference is in how the correlated yield is handled

$$sig = UnlikeFG - Norm * MixedUnlikeBG - CorrelatedUnlike$$

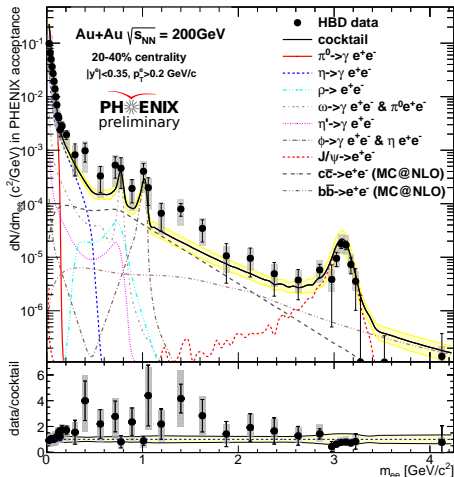
Stream A

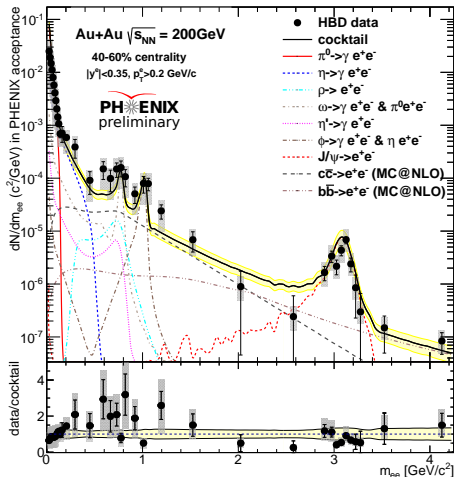
- HBD reco: Background subtracted using average charge per pad
- EID: Neural network for both Single/Double and electron identification
- Correlated dielectron background subtraction using acceptance corrected like sign

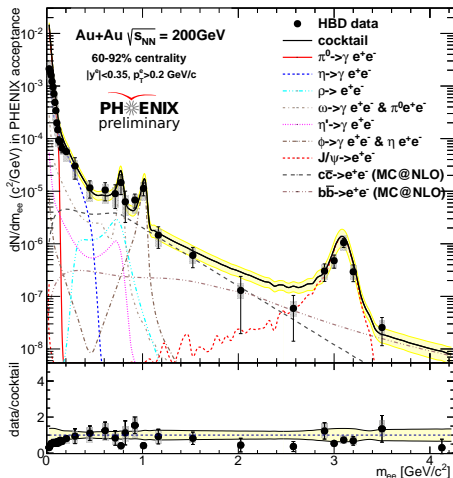
Stream B

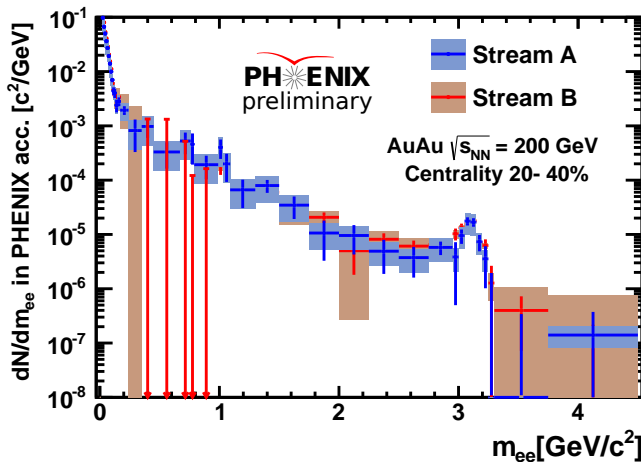
- HBD reco: Background subtracted using track projection neighborhood
- EID: Standard 1D cut for both Single/Double and electron identification
- Correlated dielectron background subtraction using MC of cross pairs and jet pairs

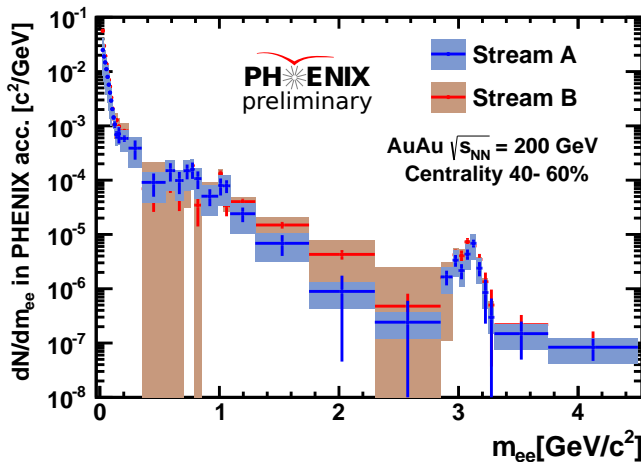
- **Stream A** is used for comparison to cocktail because the PID procedure was superior.
Stream B is used as a cross check.

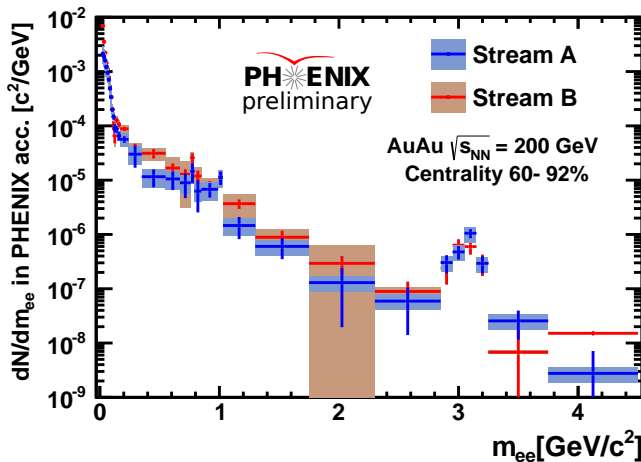


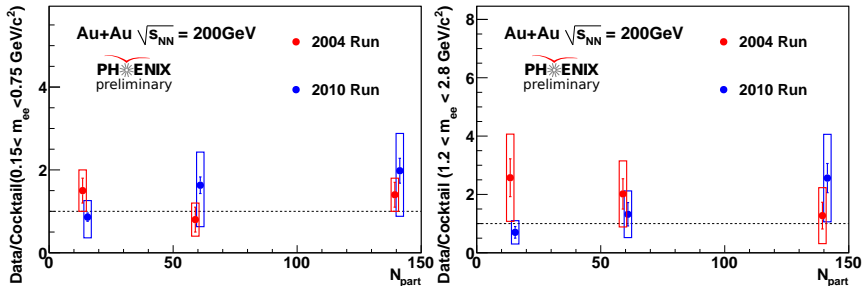




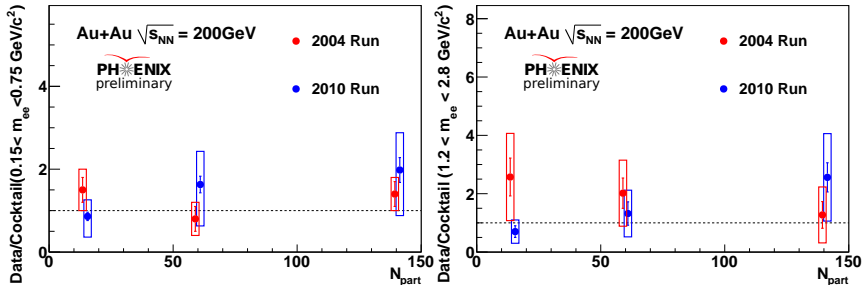








- The data/cocktail ratios comparison
 - The HBD analysis data/cocktail ratios are in agreement with the published result
 - Systematic uncertainties are dominated by conservatively estimated background subtraction errors



• Opportunities for improvement

- Extremely tight cuts and run QA used in the interest of uniformity that can be relaxed in future analyses
- The dominant contribution to the systematics comes from the background subtraction method chosen. We expect improvement in systematic uncertainties from alternate subtraction methods

- [illegible]

Backup Slides

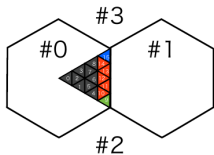
- Change in magnetic field configuration \rightarrow change in acceptance

$$\phi_{min} \leq \left(\phi - \frac{q \times k_{DC}}{p_T} \right) \leq \phi_{max}$$

$$\phi_{min} \leq \left(\phi - \frac{q \times k_{RICH}}{p_T} \right) \leq \phi_{max}$$

- where $k_{DC} = 0.058$ and $k_{RICH} = 0.127$, with $\phi_{min} = -0.55$ and $\phi_{max} = 0.957$ for one arm and $\phi_{min} = 2.185$ and $\phi_{max} = 3.686$

1. Subtract the average charge per pad on event-by-event basis for each module
2. Decide which pad to use according to projection point

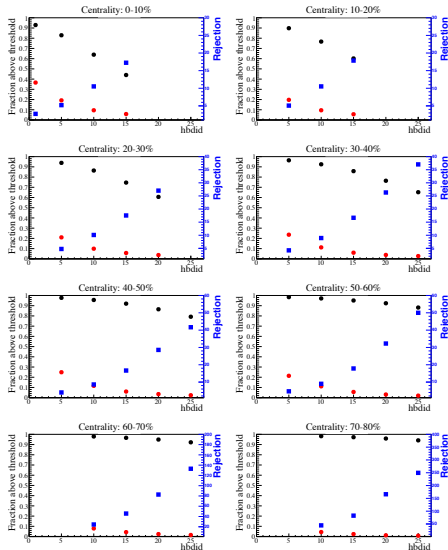


Expected cluster
 Black: #0
 Red: #0, #1
 Green: #0, #1, #2
 Blue: #0, #1, #3

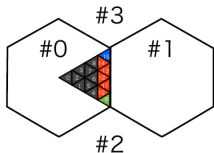
This choice will be justified in slide 10

3. Calculate the total charge in the cluster

Plots to the right: efficiency for singles and rejection for back-plane conversions for all centralities in Au+Au events.



1. Subtract the average charge per pad on event-by-event basis for each module
2. Decide which pad to use according to projection point

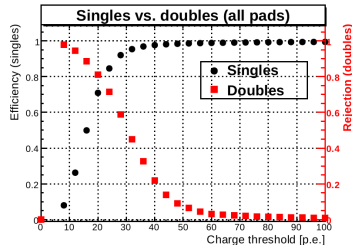
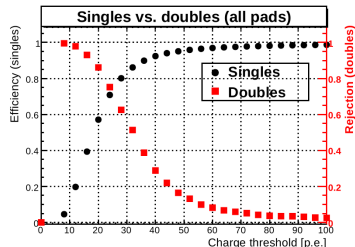


Expected cluster
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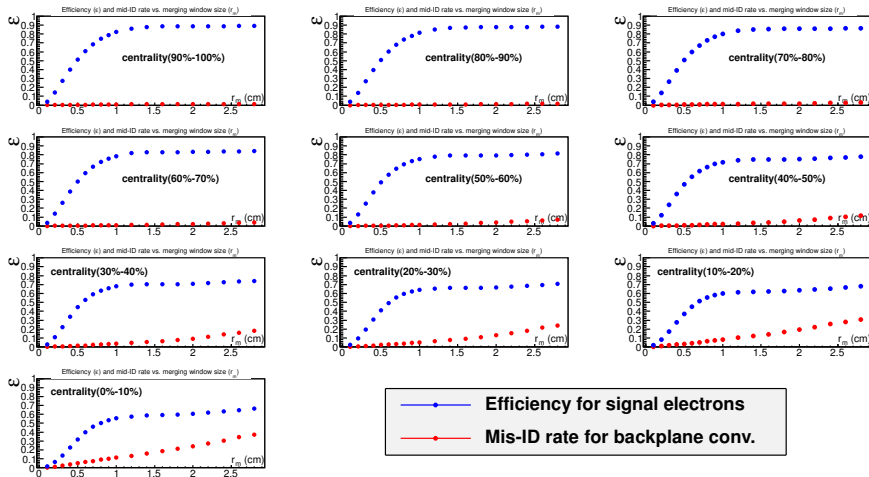
This choice will be justified in slide 10

3. Calculate the total charge in the cluster

Plots to the right: efficiency for singles and rejection for double hits for centralities 0-10% (top) and 80-90% (bottom)



- The SB reconstruction subtracts local background on triplets around track projections and merges all triplets with CG close to the track projection



- Star also has a measurement in a somewhat different acceptance
- Star also observes low mass excess compared to expectation
- However there is disagreement in the amount of the excess seen by STAR and PHENIX
- Although some of it can be attributed to acceptance difference and systematic errors, it is very desirable to have another measurement with very different systematics

